RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: $\frac{10|762,159}{1Fw0}$ Date Processed by STIC: $\frac{10|762,159}{1Fw0}$

ENTERED

RAW SEQUENCE LISTING PATENT APPLICATION US/10/762,159

DATE: 06/22/2005 TIME: 09:46:27

INPUT SET: S30675.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

```
1
                                       SEQUENCE LISTING
 2
           General Information:
 3
    (1)
          (i) APPLICANT: Boyle, William J.
 6
                         Lacey, David L.
 7
                         Calzone, Frank J.
                         Chang, Ming-Shi
8
9
         (ii) TITLE OF INVENTION: OSTEOPROTEGERIN
10
11
        (iii) NUMBER OF SEQUENCES: 168
12
13
         (iv) CORRESPONDENCE ADDRESS:
14
               (A) ADDRESSEE: Amgen Inc.
15
               (B) STREET: 1840 Dehavilland Drive
16
               (C) CITY: Thousand Oaks
17
               (D) STATE: California
18
19
               (E) COUNTRY: United States
20
               (F) ZIP: 91320
21
          (v) COMPUTER READABLE FORM:
22
23
               (A) MEDIUM TYPE: Floppy disk
24
               (B) COMPUTER: IBM PC compatible
               (C) OPERATING SÝSTEM: PC-DOS/MS-DOS
25
26
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
27
28
         (vi) CURRENT APPLICATION DATA:
               (A) APPLICATION NUMBER: US/10/762, 159
29
               (B) FILING DATE: 2004-JAN-20
30
               (C) CLASSIFICATION:
31
32
        (vii) PRIOR APPLICATION DATA:
33
               (A) APPLICATION NUMBER: US/09/132,985
34
35
               (B) FILING DATE:1998-AUG-12
36
               (A) APPLICATION NUMBER: 08/771,777
               (B) FILING DATE:
37
38
39
      (viii) ATTORNEY/AGENT INFORMATION:
40
               (A) NAME: Winter, Robert B.
               (C) REFERENCE/DOCKET NUMBER: A-378-CIP
41
42
43
44
    (2) INFORMATION FOR SEQ ID NO:1:
45
46
          (i) SEQUENCE CHARACTERISTICS:
```

RAW SEQUENCE LISTING PATENT APPLICATION US/10/762,159

DATE: 06/22/2005 TIME: 09:46:27

47.	(A) LENGTH: 36 base pairs	
48	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
49	_	
50	(D) TOPOLOGY: linear	
51		
52	(ii) MOLECULE TYPE: cDNA	
53		
54		
55		
56		
57	(xi) SEQUENCE DESCRIPTION: SEQ I	D NO.1.
	(XI) SEQUENCE DESCRIPTION: DEQ I	
58	TARGET AND ARROAD COOK COOK CANDON AND	INNT 36
59	AAAGGAAGGA AAAAAGCGGC CGCTACANNN NNN	ININ 1
60		
61	(2) INFORMATION FOR SEQ ID NO:2:	
62		
63	(i) SEQUENCE CHARACTERISTICS:	
64	(A) LENGTH: 16 base pairs	
65	(B) TYPE: nucleic acid	
66	(C) STRANDEDNESS: single	
67	(D) TOPOLOGY: linear	
	(D) TOPOLOGI. Timear	
68	(''') NOT DOIN E MUDE -DNA	• •
69	(ii) MOLECULE TYPE: cDNA	
70		
71		
72		
73		
74	(xi) SEQUENCE DESCRIPTION: SEQ I	ID NO:2:
75		
76	TCGACCCACG CGTCCG	16
77	20000000	
78	(2) INFORMATION FOR SEQ ID NO:3:	· · · · · · · · · · · · · · · · · · ·
-	(2) INFORMATION FOR BEG ID NO.3.	
79	(') CHOURNER CHARACTERICE	
80	(i) SEQUENCE CHARACTERISTICS:	·
81	(A) LENGTH: 12 base pairs	
82	(B) TYPE: nucleic acid	
83	(C) STRANDEDNESS: single	
84	(D) TOPOLOGY: linear	
85		
86	(ii) MOLECULE TYPE: cDNA	•
87		
88		
89	•	
90	() CECHENCE DECORPORANT CEC	ID NO.3.
91	(xi) SEQUENCE DESCRIPTION: SEQ	LD INO.3:
92		10
93	GGGTGCGCAG GC	12
94		
95	(2) INFORMATION FOR SEQ ID NO:4:	
96		
97	(i) SEQUENCE CHARACTERISTICS:	
98	(A) LENGTH: 18 base pairs	
99	(B) TYPE: nucleic acid	
22	(D) IIIUCICIC ACIU	•

RAW SEQUENCE LISTING PATENT APPLICATION US/10/762,159

DATE: 06/22/2005 TIME: 09:46:27

		INFULSEL SSUUTS.TUW
100	(C) STRANDEDNESS: single	
101	(D) TOPOLOGY: linear	
102		
103	(ii) MOLECULE TYPE: cDNA	
	(II) HODDCODD IIID. CDAM	
104		
105		
106		
107		
108	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
109	(111) 55251155 22551111111111111111111111	
	mama a a a coa coa coa com	18
110	TGTAAAACGA CGGCCAGT	16
111		
112	(2) INFORMATION FOR SEQ ID NO:5:	
113		
114	(i) SEQUENCE CHARACTERISTICS:	
115	(A) LENGTH: 18 base pairs	
116	(B) TYPE: nucleic acid	
117	(C) STRANDEDNESS: single	
118	(D) TOPOLOGY: linear	
119		
120	(ii) MOLECULE TYPE: cDNA	
121		
122		
123		
124	· · · · · · · · · · · · · · · · · · ·	
125	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
126	•	
127	CAGGAAACAG CTATGACC	18
128		
129	(2) INFORMATION FOR SEQ ID NO:6:	
130	(2) Infoldation for DDQ 1D no. o.	
	(i) apournan augramentantaa	
131	(i) SEQUENCE CHARACTERISTICS:	
132	(A) LENGTH: 20 base pairs	
133	(B) TYPE: nucleic acid	
134	(C) STRANDEDNESS: single	
135	(D) TOPOLOGY: linear	•
136		
137	(ii) MOLECULE TYPE: cDNA	•
	(II) FOURCODE IIIB. CDMA	
138	•	
139	•	
140		
141		
142	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
143		
144	CAATTAACCC TCACTAAAGG	20
	CHILITATOCC TOTICTIVATOC	20
145	(a) Typopyamion for dec in the	
146	(2) INFORMATION FOR SEQ ID NO:7:	
147		
148	(i) SEQUENCE CHARACTERISTICS:	
149	(A) LENGTH: 23 base pairs	
150	(B) Type: nucleic acid	
150 151	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
150 151 152		

RAW SEQUENCE LISTING PATENT APPLICATION US/10/762,159

DATE: 06/22/2005 TIME: 09:46:27

153 154 155 156	(ii) MOLECULE TYPE: cDNA	
157 158 159 160	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
161	GCATTATGAC CCAGAAACCG GAC	23
162 163	(2) INFORMATION FOR SEQ ID NO:8:	
164 165 166 167 168 169	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
171 172 173 174	(ii) MOLECULE TYPE: cDNA	
176	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
177 178	AGGTAGCGCC CTTCCTCACA TTC	23
179 180	(2) INFORMATION FOR SEQ ID NO:9:	
181 182 183 184	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 30 base pairs(B) TYPE: nucleic acid	
185 186	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
186 187 188 189 190	(D) TOPOLOGY: linear	
186 187 188 189 190 191 192 193	(D) TOPOLOGY: linear	
186 187 188 189 190 191 192 193 194	(D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA	30
186 187 188 189 190 191 192 193 194 195 196	(D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	30
186 187 188 189 190 191 192 193 194 195	(D) TOPOLOGY: linear (ii) MOLECULE TYPE: cDNA (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: GACTAGTCCC ACAATGAACA AGTGGCTGTG	30

RAW SEQUENCE LISTING PATENT APPLICATION US/10/762,159

DATE: 06/22/2005 TIME: 09:46:27

206			·	
207				
208				
209				
210		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
211				
212	ATA	GAAT	GC GGCCGCTAAA CTATGAAACA GCCCAGTGAC CATTC	45
213				
214	(2)	TNFO	RMATION FOR SEQ ID NO:11:	
215	_,			
		151	SEQUENCE CHARACTERISTICS:	
216		(1)	~	
217			(A) LENGTH: 21 base pairs	
218			(B) TYPE: nucleic acid	
219			(C) STRANDEDNESS: single	
220			(D) TOPOLOGY: linear	
221				
222		(ii)	MOLECULE TYPE: cDNA	
223		(,		
224				
225				
226				
227		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	
228				
229	GCC ⁷	CTAG	AA AGAGCTGGGA C	21
230				
231	(2)	TNFO	RMATION FOR SEQ ID NO:12:	
232	(2)	1111 01	Mail Told Told Day 15 No.11.	
		123	CHONTON OF CHARACTER CONTON	
233		(1)	SEQUENCE CHARACTERISTICS:	
234			(A) LENGTH: 21 base pairs	
235			(B) TYPE: nucleic acid	
236			(C) STRANDEDNESS: single	
237			(D) TOPOLOGY: linear	-
238				
239		(ii)	MOLECULE TYPE: cDNA	
240	-	(/	None of the second seco	
241				
242				
243				
244	•	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:12:	
245			·	
246	CGCC	GTGT	rc catttatgag c	21
247				
248	(2)	TNEO	RMATION FOR SEQ ID NO:13:	
	(2)	TIME	MARION TOK BLY ID NO.13.	
249		/ - 1	CROUDING GUADACHERICATOC.	
250		(1)	SEQUENCE CHARACTERISTICS:	
251			(A) LENGTH: 24 base pairs	
252			(B) TYPE: nucleic acid	
253			(C) STRANDEDNESS: single	
254			(D) TOPOLOGY: linear	
255				
256		(44)	MOLECULE TYPE: cDNA	
		(+ +)	MODECODE IIFE. COM	
257				
258				